

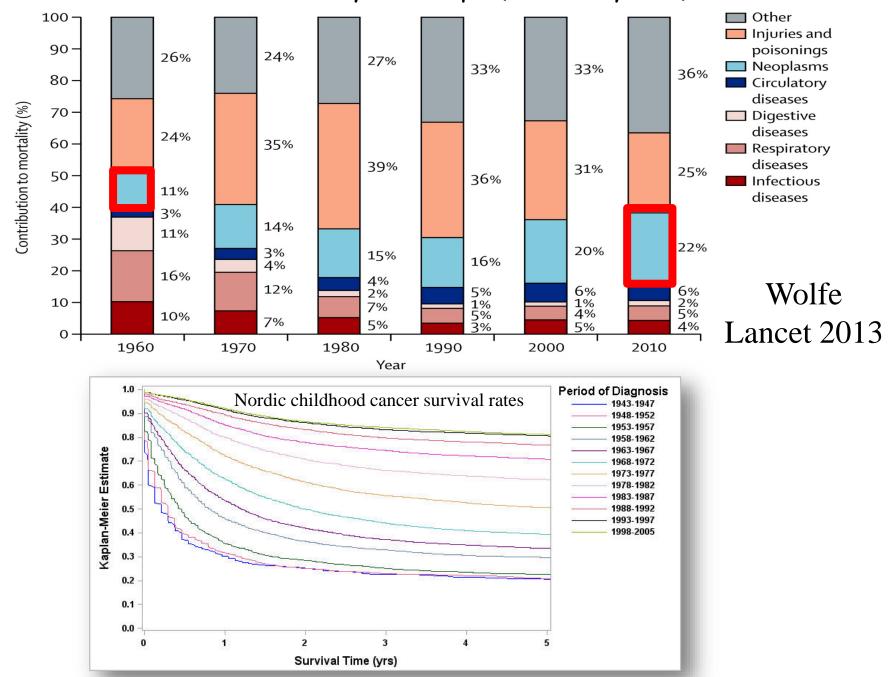
Nordic clinical trials and registries in a pediatric oncology setting

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Childhood mortality in Europe (1.0-14.9 years)







6 Countries, 6 languages

Population 25 millions 5 million children

200 with ALL per year

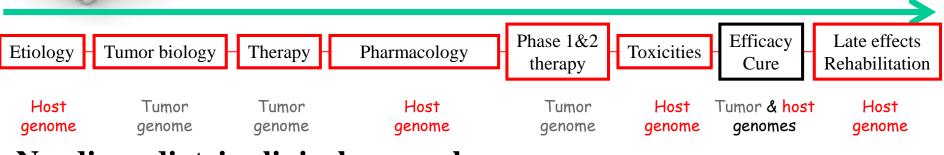
30+ ALL treatment centres

Common NOPHO protocols since 1986



Finding the genomic pieces

Patient trajectory & domains of research

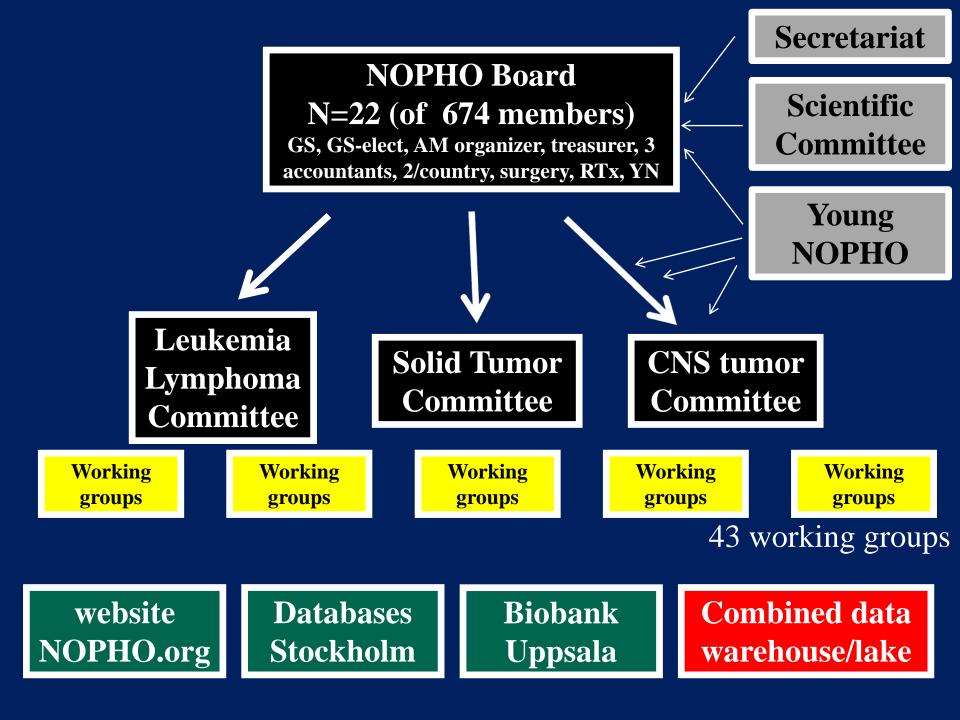


Nordic pediatric clinical research:

- Tumor biology (leukemia biobank; Uppsala)
- Dynamics of biomarkers (cytokines; pharmacology vs clinical outcome (residual leukemia; relapse; death in remission; second cancers)
- Clinical interventions: Observational (non-Rx) & Rx studies
- 1st NOPHO randomised study 1992 pharmacology of maintenance therapy: 538 pts (97% of all eligible); 10,000 samples; 30,000 treatment data sets; >20 publications

Schmiegelow, since 2009:

173 publications
50% Nordic;
+6% potentially (DK register studies)
<u>NCU supported studies:</u>
Epidemiology, pharmacology, tumor biology, clinical interventions (+/- Rx), outcome



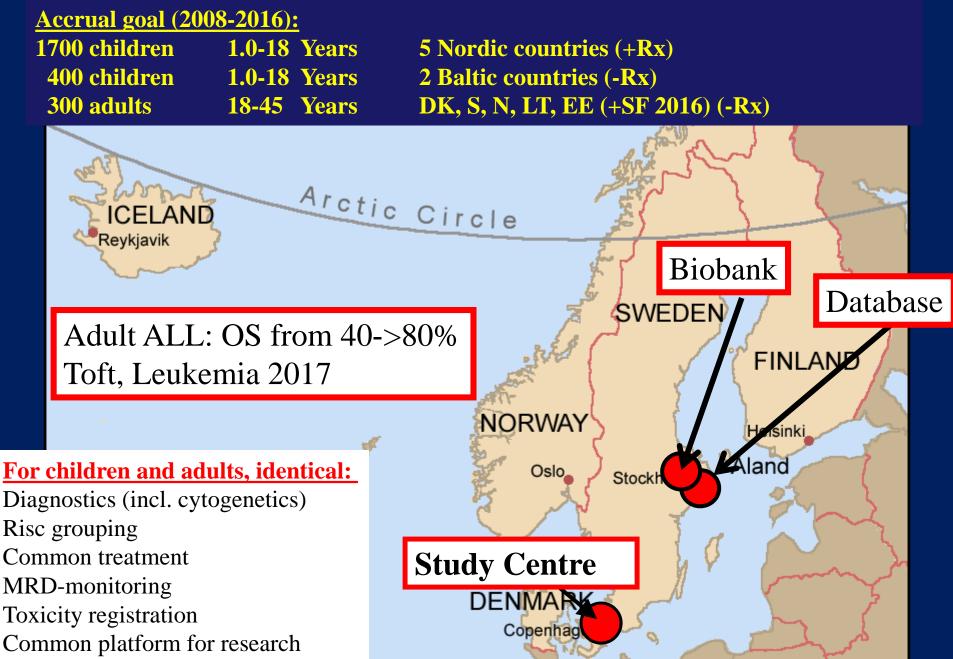
Acute Lymphoblastic Leukemia 25% of all pediatric cancers

- Rare disease
- Little interest of the pharma industry - Except immunotherapy
- Purely investigator-initiated/-driven research
 - Central Database
 - Biobank
 - Nordic prospective trials / studies

Nordic Clinical ALL Trial Challenges

- 5 (7) countries, 5 (7) languages
- 5 (7) national authorities Medicines Agencies
- 5 (7) National Data protection agencies
- 5 (7) Ethical Boards
- 5 (7) interpretations of the EU Directive and 5 (7) sets of ethical rules
- Strategy and funding for GCP monitoring in Investigator Initiated trials

NOPHO ALL-2008



Strategy of monitoring and registration



Biggest challenge in the Nordic Pediatric Oncology setting: Ethical applications – not registries

• One entry

Or perhaps

• VHP- like ethical application

Nordic Cooperation

Strategy for the trial

- Simple, on-line dataregistration, including SAEs, Death and SUSAR's
- Exclusion of known AE's
- Continuous monitoring of entered data by the study centre. Errors are picked up within a short period.
- Nordic GCP network
- Help-desk

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Strategy for the trial

- Simple, on-line dataregistration, including SAEs
- Exclusion of known AE's
- Continuous monitoring of registered data by the study centre. Errors are picked up within a short period.
- Nordic GCP network
- Help-desk

AEs not to be reported

- a number of toxicities are so well-known and frequent during therapy that they will not be reported. These includes:
- For the 6MP increment study, the following will not be AE-reported:
- Since leukopenia is the target toxicity (monitoring parameter), this side-effect will not be regarded as a SAE. This also includes febrile neutropenia leading to hospitalisation or prolongation of ongoing hospitalisation if the patients condition otherwise is good with no signs of septic shock.
- Since <u>thrombocytopenia</u> is the target toxicity (monitoring parameter) this side-effect will not be regarded as a SAE.
- A rise in aminotransferases with normal liver function tests (i.e. bilirubin and INR (or coagulation factor 2-7-10) is a well-known side effect of HD-MTX and 6MP and will not be regarded as a SAE, unless in combination with 19.3.1.8.
- \bullet **A rise in bilirubin** to less than 5x UNL.
- A <u>fall in coagulation factors</u>, unless in combination with 19.3.1.8.
- Less than a grade 4 rise in amylase (>5x UNL, if measured) will not be reported.
- Kidney dysfunction is a well-known side effect of HD-MTX and will not be regarded as SAE unless it requires dialysis or leads to a permanent kidney dysfunction with s-creatinine >UNL.
- Stomatitis and dyspepsia with or without liver toxicity are a well-known side effects of HD-MTX and will not be regarded as SAE.
- **Infection/fever** leading to hospitalisation or prolongation of existing hospitalisation.

Nordic Cooperation

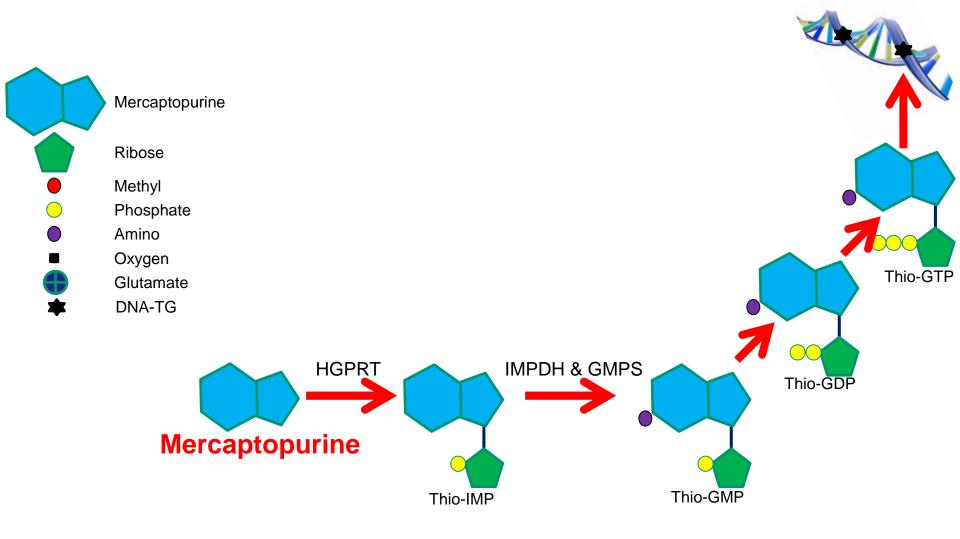
Compliance

- <u>Quarterly registrations:</u> 33 out of 34 centres (97%) register within 1 months (both adult and child centers)
- >99 % of eligible patients participate in the common treatment protocol (2¹/₂ years)
- 80-85% participate in randomizations

Acute Lymphoblastic Leukemia 25% of all pediatric cancers

- Rare disease
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 - Central Database
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 - Nordic prospective trials / studies
- From 2019: NOPHO →

ALLtogether (14 European countries)



Schmiegelow et al. JPHO 2014 (review)

NOPHO ALL2008 Maintenance therapy study

• 1016 non-HR ALL patients were eligible

- 7 no family consent
- 101 no samples taken
- 918 included (89% of all eligible)
 - 526 MRD-positive day 29
 - 390 MRD-negative day 29
 - 2 no MRD status
- 5y-EFS: 92.4% (40 relapses)
- Standard risk (N=549; 60%)
 - BCP with MRD <0.1% day 29</p>
 - No CNS3
 - No i21amp
 - No t(1;19)
 - No dic(9;10)

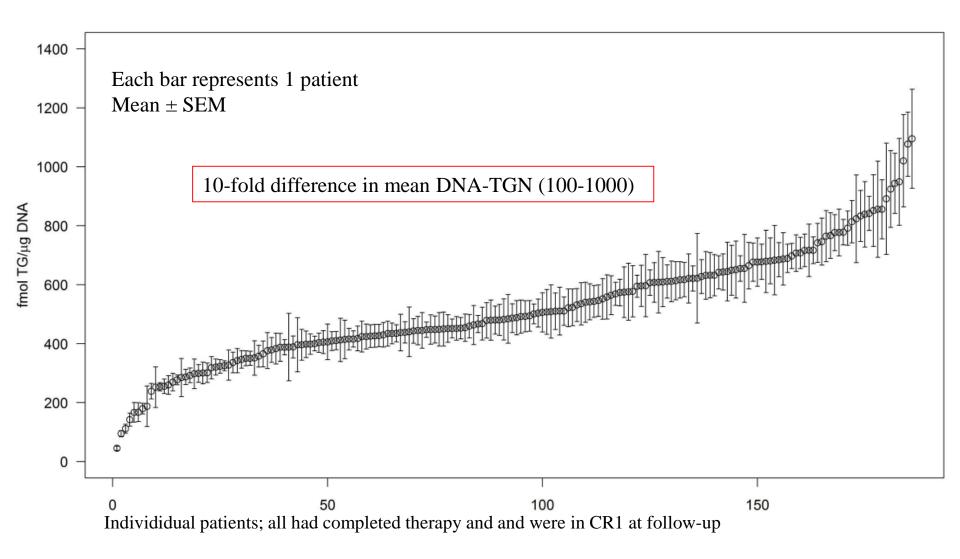
	Patients (n=918)	
Age at diagnosis (years)	4.2 (2.9–7.3)	
Sex		
Male	489 (53%)	
Female	429 (47%)	
White blood cell count at diagnosis (× 10° cells per L)	9.2 (4.3-30.9)	
Risk group		
Standard	549 (60%)	
Intermediate	369 (40%)	
Immunophenotype		
B-precursor leukaemia	854 (93%)	
T-cell leukaemia	64 (7%)	

Data are n (%) or median (IQR).

- 346 in CR1 at end of therapy & >5 samples in 6MP/MTX maintenance
 - Pharmacological modelling

186 patients with ≥10 DNA-TGN measurements during last 1.5 years of maintenance NOPHO ALL2008 maintenance therapy study

Patients can be classified according to their DNA-TGN



Risk of relapse by DNA-TGN NOPHO ALL-2008 918 non-HR patients reaching start of maintenance therapy Measurements per patient (in MT-2, only 6MP/MTX); median N=9 (1-56) >10,000 blood samples

	Positive MRD day 29 n = 526, 31 relapses			Negative MRD day 29 n = 390, 9 relapses			
	Relapse specific HR	95% CI	p-value	Relapse specific HR	95% CI	p-value	
DNA-TGN per 100 ^a	0.723	0.572–0.913	0.0065	1.010	0.733–1.391	0.95	
Age at diagnosis	1.118	1.037–1.205	0.0035	1.073	0.923–1.247	0.36	
Female sex	1.036	0.511–2.100	0.92	0.613	0.149–2.524	0.49	
WBC at Dx per 10x10 ⁹ /L	1.001	0.998–1.005	0.56	1.005	1.007–1.097	0.022	
4							

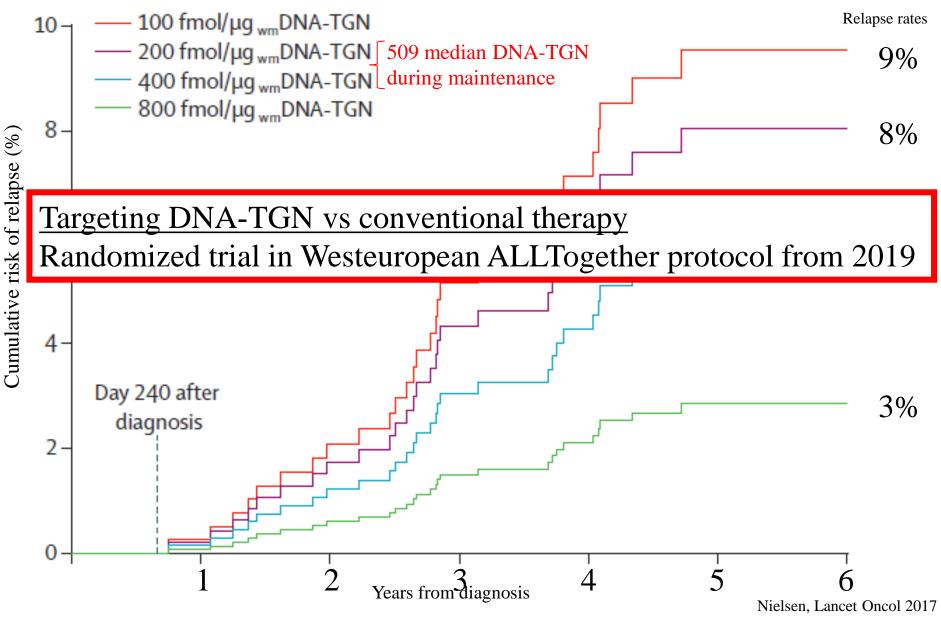
^a Time-dependent mDNA-TGN level re-calculated at each time of event

28% reduction in relapse hazard risk per increment of DNA-TGN of 100 fmol/µg DNA

Simulated relapse curves for boy, 5.0 yrs, WBC 9.6 at Dx, in CR1 d240.

28 relapses in 494 day 29 MRD-positive children Min 1 DNA-TGN measurement before day 240 - then fixed

Proportional hazards model



STAGING Sequencing Tumor And Germline DNA – Implications and National Guidelines

STAGING Sequencing of Tumor And Germline DNA – Implications and National Guidelines

DAY 1	WEEK 1-2	WEEK 3-4	WEEK 5-6	
PEDIATRIC CANCER DIAGNOSIS	WRITTEN MATERIAL + SHORT ORAL PRESENTATION	GENETIC COUNSELING + DEBRIEFING	INTERVIEW + BLOOD SAMPLE	WHOLE GENOME SEQUENCING

STAGING

Sequencing Tumor And Germline DNA – Implications and National Guidelines

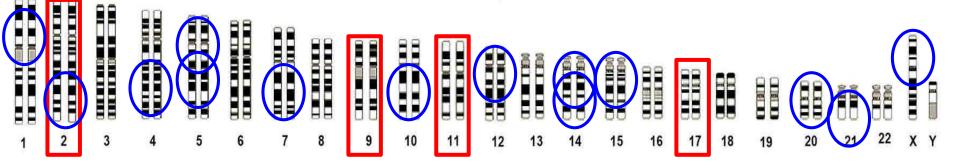
STAGING – Norway

WGS/WES/tumor RNAseq ++, BCF-Sweden

Copenhagen-Lund collaboration Copenhagen-Vilnius collaboration

The "competition" is scientific & political & financial

A look into germline DNA



ATM mutation; ~50-100x riskof lymphoma/ALL

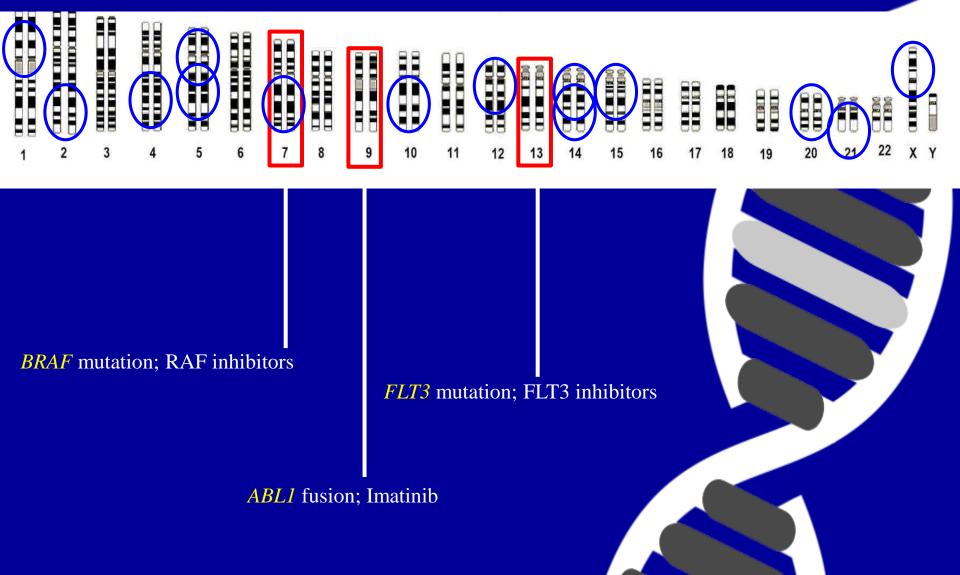
Biallel MMR mutations; ~100% absolute risk of cancer <18y

cancer, 1/3 <18y

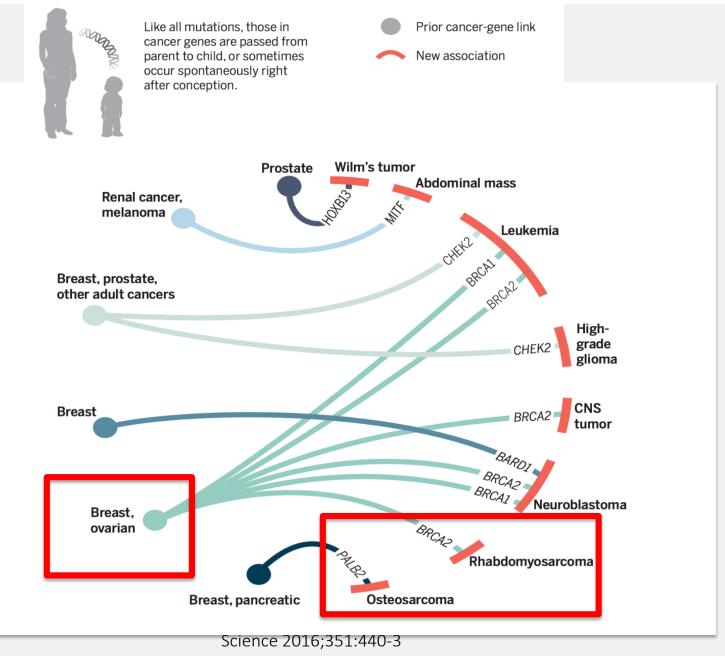
TP53 mutation: 90-100% life time risk of

PAX5 mutation: Markedly increased risk of ALL

A look into germline DNA



Germline mutations in adult and childhood cancer



Whole genome sequencing and data analysis

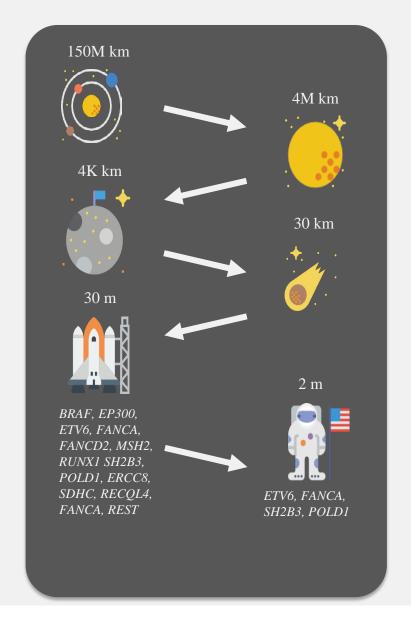
100 billion bp reads *Alignment with "standard" genome*3 billion bp (genome) *Variant calling*3 million variants *In or around an exome*20.000 variants *Rare <1%*2.000 variants *216 cancer predisposition genes*20 variants *Depth, quality, impact, frequency, in silico prediction, conservation (evolution), cancer*

type and mutations, family history, litterature curation

0-4 variants

Multidisciplinary team conference (weekly,10-20% af alle pts)

No standards exist internationally for the bioinformatics pipeline or for reporting Some consensus on follow-up



"Pediatric cancer families' participation in whole genome sequencing research in Denmark: parent perspectives"

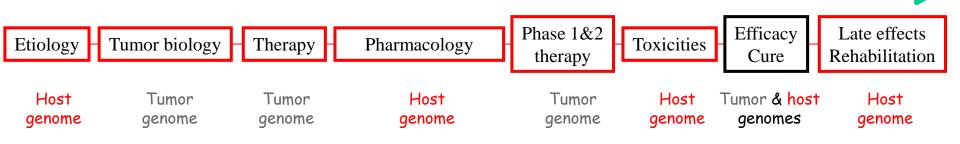
STUDY FINDINGS: (15 families (30 parents), written info 2-28 days, genetic counselling 7-42 days after diagnosis)

- When is the right time? Most had no objections to being approached / counselled within 4 weeks from diagnosis.
- A few parents find it too early.
- **Why has this happened?** Parents have many questions about cancer risk including genetic
- **Making the right decision** Parents have concerns regards secondary findings and expressed that they *may* end up regretting consent. Many families had very in-depth discussions about which findings to have reported back, at times with discordant views between parents.

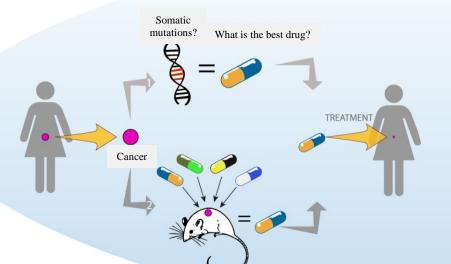


Finding the genomic pieces Personalized medicine

Patient trajectory & domains of research



PERSONALIZED CANCER MEDICINE



Personalized medicine in ped. oncology:

- Cancer predisposition
- Tumor -omics (diagnosis/prognosis)
- Phase 1 & phase 2 trials
- Therapeutic drug monitoring
- Toxicities (treatment to the limit of toxicity)
- Genotype-phenotype vs phenotype-phenotype

The funding: Money and Politics

- Nordic studies are especially important
- if the Nordic group has a unique *international role*
- if they strengthen Nordic *collaboration* not just data provision
 - e.g. shared PhD-students required
- if a Nordic setting is needed (for study power), since the questions are unique for the Nordic *"culture"* or future Nordic *health care*